



STIC Search Report

Biotech-Chem Library

STIC Database Tracking Number: 142006

**TO: Terra Gibbs
Location: 2d10 / 2c18
Tuesday, January 25, 2005
Art Unit: 1635
Phone: 272-0758
Serial Number: 10 / 005337**

**From: Jan Delaval
Location: Biotech-Chem Library
Rem 1a51
Phone: 272-2504

jan.delaval@uspto.gov**

Search Notes

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SEARCH REQUEST FORM

Scientific and Technical Information Center

Requester's Full Name: _____ Examiner #: _____ Date: _____
 Art Unit: _____ Phone Number 30 _____ Serial Number: _____
 Mail Box and Bldg/Room Location: _____ Results Format Preferred (circle): PAPER DISK E-MAIL

If more than one search is submitted, please prioritize searches in order of need.

Please provide a detailed statement of the search topic, and describe as specifically as possible the subject matter to be searched. Include the elected species or structures, keywords, synonyms, acronyms, and registry numbers, and combine with the concept or utility of the invention. Define any terms that may have a special meaning. Give examples or relevant citations, authors, etc, if known. Please attach a copy of the cover sheet, pertinent claims, and abstract.

Title of invention: _____

Inventors (please provide full names): _____

Earliest Priority Filing Date: _____

**For Sequence Searches Only* Please include all pertinent information (parent, child, divisional, or issued patent numbers) along with the appropriate serial number.*

STAFF USE ONLY

Searcher: Jan
 Searcher Phone #: 22504
 Searcher Location: _____
 Date Searcher Picked Up: 1/25/05
 Date Completed: 1/25/05
 Searcher Prep + Review Time: _____
 Clerical Prep Time: 10
 Online Time: + 45

Type of Search

NA Sequence (#) ☒
 AA Sequence (#) _____
 Structure (#) _____
 Bibliographic _____
 Litigation _____
 Fulltext _____
 Patent Family _____
 Other _____

Vendors and cost where applicable

STN _____
 Dialog _____
 Questel/Orbit _____
 Dr.Link _____
 Lexis/Nexis _____
 Sequence Systems ☒
 WWW/Internet _____
 Other (specify) _____

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142006

Delaval, Jan

From: Gibbs, Terra
Sent: Tuesday, January 25, 2005 2:16 PM
To: Delaval, Jan
Subject: RE: 10/005337

The Accession number is AF041847.

-----Original Message-----

From: Delaval, Jan
Sent: Tuesday, January 25, 2005 8:08 AM
To: Gibbs, Terra
Subject: 10/005337

Terra -

I am processing your search request for 10 / 005337.

You have requested a comparison between seq id no 2 and af04184.

I have not been able to locate this accession number in any of our in-house databases; I could not locate this number at NCBI.

Please verify the accession number.

Thanks.

Jan Delaval, 22504

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142006

STIC-Biotech/ChemLib

From: Gibbs, Terra
Sent: Tuesday, January 18, 2005 4:42 PM
To: STIC-Biotech/ChemLib
Subject: RE: Sequence comparison

I submitted this request, but put the wrong Accession Number,

I requested a search for SEQ ID NO: 2 of USSN 10/005,337. Accession number **AF04184** came up as a good piece of art.

However, I need Accession Number **AF04184** to be at least 80% identical to SEQ ID NO:2 of USSN 10/005,337 or as close as possible.

Can I please have a comparison between these two sequences, with the similarity being at least 80%?

Thank You.

-----Original Message-----

From: Gibbs, Terra
Sent: Friday, January 07, 2005 4:24 PM
To: STIC-Biotech/ChemLib
Subject: Sequence comparison

I requested a search for SEQ ID NO: 2 of USSN 10/005,337. Accession number AF131884 came up as a good piece of art.

However, I need Accession Number AF131884 to be at least 80% identical to SEQ ID NO:2 of USSN 10/005,337.

Can I please have a comparison between these two sequences, with the similarity being at least 80%?

Thank You.

*Terra Cotta Gibbs, Ph.D.
Art Unit 1635
Remsen Building 2D10
Mailbox 2C18
571-272-0758*

STAFF USE ONLY

Searcher: _____
Searcher Phone: 2- _____
Date Searcher Picked up: _____
Date Completed: _____
Searcher Prep/Rev. Time: _____
Online Time: _____

Type of Search

NA Sequence: # _____
AA Sequence: # _____
Structure: # _____
Bibliographic: _____
Litigation: _____
Patent Family: _____
Other: _____

Vendors and cost where applicable

STN: _____
DIALOG: _____
QUESTEL/ORBIT: _____
LEXIS/NEXIS: _____
SEQUENCE SYSTEM: _____
WWW/Internet: _____
Other(Specify): _____

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STIC-Biotech/ChemLib

1412004

From: Gibbs, Terra
Sent: Friday, January 07, 2005 4:24 PM
To: STIC-Biotech/ChemLib
Subject: Sequence comparison

I requested a search for SEQ ID NO: 2 of USSN 10/005,337. Accession number AF131884 came up as a good piece of art.
However, I need Accession Number AF131884 to be at least 80% identical to SEQ ID NO:2 of USSN 10/005,337.
Can I please have a comparison between these two sequences, with the similarity being at least 80%?
Thank You.

Terra Cotta Gibbs, Ph.D.
Art Unit 1635
Remsen Building 2D10
Mailbox 2C18
571-272-0758

RECEIVED
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(STIC)

STAFF USE ONLY

Searcher: _____
Searcher Phone: 2- _____
Date Searcher Picked up: _____
Date Completed: _____
Searcher Prep/Rev. Time: _____
Online Time: _____

Type of Search

NA Sequence: # _____
AA Sequence: # _____
Structure: # _____
Bibliographic: _____
Litigation: _____
Patent Family: _____
Other: _____

Vendors and cost where applicable

STN: _____
DIALOG: _____
QUESTEL/ORBIS: _____
LEXIS/NEXIS: _____
SEQUENCE SYSTEM: _____
WWW/Internet: _____
Other(Specify): _____

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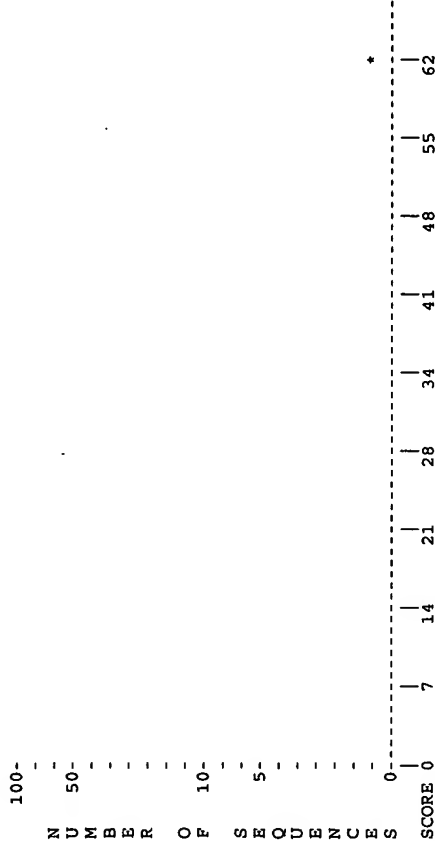
> O <
O | O IntelliGenetics
> O <

FastDB - Fast Pairwise Comparison of Sequences
Release 5.4

Results file seq2-af041847.res made by jdelaval on Tue 25 Jan 105 14:35:43-PST.

Query sequence being compared:US-10-005-337A-2 (1-2074)
Number of sequences searched: 1
Number of scores above cutoff: 1

Results of the initial comparison of US-10-005-337A-2 (1-2074) with:
File : af041847.seq



PARAMETERS

Similarity matrix Unitary K-tuple 4
Mismatch penalty 1 Joining penalty 30
Gap penalty 1.00 Window size 32
Gap size penalty 0.33
Cutoff score 0
Randomization group 0

SEARCH STATISTICS

Scores: Mean 62 Median 0 Standard Deviation 0.00
Times: CPU 00:00:00.00 Total Elapsed 00:00:00.00
Number of residues: 1026
Number of sequences searched: 1
Number of scores above cutoff: 1

The scores below are sorted by initial score.
Significance is calculated based on initial score.
A 100% identical sequence to the query sequence was not found.
The list of best scores is:

Sequence Name Description Length Score Score Init. Opt.

1. af041847 TOIG of: af041847 check: 453 1026 62 434 0.00 0
1. US-10-005-337A-2 (1-2074)
af041847 TOIG of: af041847 check: 4536 from: 1 to: 1026
TOIG of: af041847 check: 4536 from: 1 to: 1026
LOCUS AF041847 1026 bp mRNA linear ROD 21-FEB-1998
DEFINITION Mus musculus cardiac ankyrin repeat protein MCRP mRNA, complete
cds.
ACCESSION AF041847
VERSION AF041847.1 GI:2905615
KEYWORDS
SOURCE Mus musculus (house mouse)
ORGANISM
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mus.
1 (bases 1 to 1026)
REFERENCE
AUTHORS Zou,Y., Evans,S., Chen,J., Kuo,H.C., Harvey,R.P. and Chien,K.R.
TITLE CARP, a cardiac ankyrin repeat protein, is downstream in the Nkx2-5
homeobox gene pathway
JOURNAL Development 124 (4), 793-804 (1997)
MEDLINE 97195688
PUBMED 9043061
REFERENCE 2 (bases 1 to 1026)
AUTHORS Chen,J. and Chien,K.R.
TITLE Direct Submission
JOURNAL Submitted (07-JAN-1998) Medicine, UCSD, 9500 Gilman Dr, La Jolla,
CA 92093, USA
FEATURES
source Location/Qualifiers
1..1026
/organism="Mus musculus"
/mol_type="mRNA"
/db_xref="taxon:10090"
/tissue_type="heart"
15..974
/codon_start=1
/product="cardiac ankyrin repeat protein MCRP"
/protein_id="AAC03533.1"
/db_xref="GI:2905616"
/translation="MMVLRVBELVTGKNSNGAAGFLPGFRNGEYAAVALEKQED
LKTIPANSVKGEORSEKLEKLEQSKLENLEDTIIVOLKKRKYKKT
KVPVVKPEPEINTEPVDVPRFLKALENKLPVVEKLVSDKNSPDVCEYKRTALHRA
CLEGHLAIVEKLEAGQIEPRDMLESTAIHWACRGNADVLKLLNKGAKISARDKL
LSTALHVAVRTGHYECABHLIACEDLNADREGDTPLHDAVRLNRYKMIRLMTFGA
DLKVKNCAGKTPMDLVLHWQSGTKAIFDSPKENAYKNSRIATF"
ORIGIN
AF041847 Length: 1026 January 25, 2005 14:32 Type: N Check: 4536 ..
Initial Score = 62 Optimized Score = 434 Significance = 0.00
Residue Identity = 48% Matches = 549 Mismatches = 407
Gaps = 168 Conservative Substitutions = 0
10 20 30 40 50 60 70
CTGCAGCAAGTTACTTAATGTTTTTTCCTCAGCATCCTCTCTGTAAATAGAGAGCATTAGTCTGCTCCAA
80 90 100 110 120 130 140
CTTCGAGGCATGGACAGCTCTGGGATTTTCATATCCAGACCCCTTAAACATCCACAGTCTTCCCCCAAC
150 160 170 180 190 200 210
ACTTCTCCTCCTTAATACCTCCCTCAGTTGGGTGGCTGAGCAACAAAGGCATACGAAATGGTAGAAAA
220 230 240 250 260 270 280
GTGTCCATGACTACTTCTGACTTAGATGAAGAGACCAATGAAATAGTAATGACTCTGTTTGTCTCAGCAGG
290 300 310 320 330 340 350 360
ACATATCTAAATAGGAGCTATACAAAGAGATTAGCATGGACTCTGTGCAAGATGACACACAATTTGT

IntelliGenetics

FastDB - Fast Pairwise Comparison of Sequences
Release 5.4

Results file us-10-005-337a-2-inv.res made by jdelaval on Tue 25 Jan 105 14:36:34-PST.

Query sequence being compared: US-10-005-337A-2' (1-2074)
 Number of sequences searched: 1
 Number of scores above cutoff: 1

Results of the initial comparison of US-10-005-337A-2' (1-2074) with:
File : af041847.seq

GRADE	NUMBER OF STUDENTS
A	28
B	25
C	22
D	19
E	16
F	12

Sequence Name	Description	Length	Score	Init. Opt.
1. af041847	TOIG of: af041847	453	1026	28 423 0.00 0

1. US-10-005-337A-2' (1-2074)
af041847 TOIG Of: af041847 check: 4536 from: 1 to: 1026

TOIG of: af041847 check: 4536 from: 1 to: 1026

LOCUS AF041847 1026 bp mRNA ROD 21-FEB-1998
DEFINITION Mus musculus cardiac ankyrin repeat protein MCARP mRNA, complete cds.

ACCESSION	AF041847
VERSION	AF041847.1
KEYWORDS	GI:2905615
SOURCE	Mus musculus (house mouse)
ORGANISM	Mus musculus
	Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Mus.
REFERENCE	1 (bases 1 to 1026)
AUTHORS	Zou, X., Evans, S., Chen, J., Kuo, H.C., Harvey, R.P. and Chien, K.R.
TITLE	CARP, a cardiac ankyrin repeat protein, is downstream in the Nkx2-5 homeobox gene pathway
JOURNAL	Development 124 (4): 793-804 (1997)

ORIGIN

AF041847 Length: 1026 January 25, 2005 14:32 Type: N Check: 4536 ..

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Initial Score      = 28      Optimized Score = 423      Significance = 0.00
Residue Identity  = 46%     Matches       = 513     Mismatches  = 453
Gaps              = 135     Conservative Substitutions = 0

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GTGGCTGAAGGAGCTCTGTATGTTTCTGTCTGTGGAGGAATCCCTGGAGTTGGCCCTGCTGGGCCCTC

80	90	100	110	120	130	140
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	150	160	170	180	190	200	210
150	150	150	150	150	150	150	150
160	150	160	150	150	150	150	150
170	150	150	170	150	150	150	150
180	150	150	150	180	150	150	150
190	150	150	150	150	190	150	150
200	150	150	150	150	150	200	150
210	150	150	150	150	150	150	210

[illegible]

90 300 310 320 330 340 350 360

370 380 390 400 410 420 430
ACCACATCACTGCCCTTTTCTTTCTTTGTCAGCTTTTCATATGACTACTCTATCAAGAAATGTAGATGCCCTAC
440 450 460 470 480 490 500
ATATCAACCCCGAGTAATATCTTTCTGTATAGCAGACTTATCAACACATTCACCTTAGGGGAACTTGTCCCA
510 520 530 540 550 560 570
GGACATCTATTCCCTGGGTAAACAGCCTTGAGGGGAAGGATCTGGGCCCTAAAGSCACTTGTCTATATCTG
580 590 600 610 620 630 640
TTTGGAAATCTTTGGAGTGAATGTCTGGGCTAAATAGGCCCATGACATGCCACTTTACCATCATATATAGAC
650 660 670 680 690 700 710 720
ATTGAAGCATTCAGTAATGCACTTTCTTTTATGATGTCACTGGCCATTAGCACATGACATGCTTTACCGTG
730 740 750 760 770 780 790
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800 810 820 830 840 850 860
GAAACCAGCTATCAGCCGGGTGCACTGGCTTACACCTGTAATCCCAACATTTTGGGAGGCAGACAGGAAG
870 880 890 900 910 X 920 930
ATTGCTTAGCTTAGGAGTTCATACTAGCTGGCAACATGGCAAAATCCCATCT-CTACCAAA--ATACT
TCACGGCTGCCAACATGATGTACT
X 10 20
940 950 960 970 980 990 1000
-ATA-TATA-TATATGAVAAATAGACCAAGATATAGCAATGGCAGACTCTGATTAGGTCAATGAGCTG
1010 1020 1030 1040 1050 1060 1070
AAGTCTCAAAATGCTGTGGGAAAGAAATCACTGTGGAGCTTGTCAAGAAATACAGA-TCCCTGGCCCAAC
1080 1090 1100 1110 1120 1130 1140
TCAGGGAATCTGAGTTGATTTATGTTAA--CTCGTAGGAATCATCTTGAATTTAAAGCAACCCAGA--TG
1150 1160 1170 1180 1190 1200
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1210 1220 1230 1240 1250 1260 1270
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1280 1290 1300 1310 1320 1330 1340 1350
GTTTCAGGAAACCTGGGTTTMTSAGTAAAGA CAGGAGAAAGATGGGAGAAAGGAAATGTGAATGGA
1360 1370 1380 1390 1400 1410 1420
GGGAGTGTAAATAGGAAGTATGTTGACCTTTGTG-TGTCTATTTCTCCCAAACTCAC-CCATCAAC
1430 1440 1450 1460 1470 1480
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CGAGCATGCTTAGA-AGGACATCTGGCGATCGTGGAGAAATTAAT-GGAGGCTGGAG-CCGAGTTGAATTC

440 450 460 470 480 490 500
ATATCAACCCCGAGTAATATCTTTCTGTATAGCAGACTTATCAACACATTCACCTTAGGGGAACTTGTCCCA
510 520 530 540 550 560 570
GGACATCTATTCCCTGGGTAAACAGCCTTGAGGGGAAGGATCTGGGCCCTAAAGSCACTTGTCTATATCTG
580 590 600 610 620 630 640
TTTGGAAATCTTTGGAGTGAATGTCTGGGCTAAATAGGCCCATGACATGCCACTTTACCATCATATATAGAC
650 660 670 680 690 700 710 720
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730 740 750 760 770 780 790
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870 880 890 900 910 X 920 930
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TCACGGCTGCCAACATGATGTACT
X 10 20
940 950 960 970 980 990 1000
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1010 1020 1030 1040 1050 1060 1070
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1080 1090 1100 1110 1120 1130 1140
TCAGGGAATCTGAGTTGATTTATGTTAA--CTCGTAGGAATCATCTTGAATTTAAAGCAACCCAGA--TG
1150 1160 1170 1180 1190 1200
ATTCTGATGCATCAAGATTGAGACACGCTGGTGGAGGATA-ATGCACCTTTTTCCTGGGAAGGTTG
1210 1220 1230 1240 1250 1260 1270
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1280 1290 1300 1310 1320 1330 1340 1350
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1360 1370 1380 1390 1400 1410 1420
GGGAGTGTAAATAGGAAGTATGTTGACCTTTGTG-TGTCTATTTCTCCCAAACTCAC-CCATCAAC
1430 1440 1450 1460 1470 1480
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CGAGCATGCTTAGA-AGGACATCTGGCGATCGTGGAGAAATTAAT-GGAGGCTGGAG-CCGAGTTGAATTC

490 500 510 520 530 540 550
1490 1500 1510 1520 1530 1540 1550
TGAGA-AGAC-----ATCGACTCCCAACCCCAAGACAAACATTGACAAAGAGGACCTTCCCAGTACCTCT
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560 570 580 590 600 610 620
1560 1570 1580 1590 1600 1610
G-TATCATATGTTCC--GGTC--TCCTA---ATATTCTATCTTA--AAGGGGTAAATAAG-GAATCTC
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630 640 650 660 670 680 690
1620 1630 1640 1650 1660 1670 1680
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1690 1700 1710 1720 1730 1740 1750
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770 780 790 800 810 820 830
1760 1770 1780 1790 1800 1810 1820
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840 850 860 870 880 890
1830 1840 1850 1860 1870 1880
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CAGAGTGAACCCAAAGCAATATTCGACAGCCCCCAAGAGAAATGCCTACAAGAACTCTCGCATAGCTACATTC
900 910 920 930 940 950 960 970
1890 1900 1910 1920 1930 1940 1950 X
TGTTCCAGGCTGACCCAAACCTGAGGGAGGTATAGGAGGAGAACTGTTGGGGAAGGACTGTGGGATGTT
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980 990 1000 1010 1020 X
1960 1970 1980 1990 2000 2010 2020
TAAGGCTCTTGGATATGAATCCCAAGCTGTCCATGCCCTGAAAGTTGGACCAAGACTAATGCTCTCATTT
2030 2040 2050 2060 2070
TACAGAGAGGATGCTGAGGCCAAACAAATTAAGTAACCTTGCTGCAG

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